

# Resource Summary Report

Generated by [kravitz2](#) on Aug 2, 2026

## MutPred Splice

RRID:SCR\_000594

Type: Tool

### Proper Citation

MutPred Splice (RRID:SCR\_000594)

### Resource Information

**URL:** <http://mutdb.org/mutpredsplice/>

**Proper Citation:** MutPred Splice (RRID:SCR\_000594)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 6,2023. Tool for identifying coding region variants which disrupt pre-mRNA splicing and the underlying mechanism.

**Abbreviations:** MutPred Splice

**Resource Type:** analysis service resource, service resource, production service resource

**Defining Citation:** [PMID:24451234](#)

**Keywords:** exonic variant, splicing, mrna, next generation sequencing, coding region substitution

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** MutPred Splice

**Resource ID:** SCR\_000594

**Alternate IDs:** OMICS\_02257

**Record Creation Time:** 20220129T080202+0000

**Record Last Update:** 20260801T190830+0000

### Ratings and Alerts

No rating or validation information has been found for MutPred Splice.

No alerts have been found for MutPred Splice.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

Mort M, et al. (2014) MutPred Splice: machine learning-based prediction of exonic variants that disrupt splicing. Genome biology, 15(1), R19.